

Serial No. 09/803,736

Examiner: Einsmann

Amendments to the Claims

This listing of claims will replace all prior versions, and listings, of claims in the application:

Listing of Claims:

Claims 1 (currently amended) A method of identifying a region of genomic DNA associated with a phenotypic trait of interest comprising:

- (A) screening a mapping population of *Arabidopsis thaliana* plants to determine the linkage of said phenotypic trait with a collection of polymorphisms, wherein at least one of the polymorphisms is selected from Table A;
- (B) calculating the linkage of each of said polymorphism to said phenotypic trait and
identifying said region of genomic DNA associated with a phenotypic trait based on linkage of said trait to one or more of said polymorphisms

and wherein said collection of polymorphisms is designed from features of polymorphisms listed in Table A including Single Nucleotide Polymorphism 471736 and recorded on computer readable medium.

Claims 2-29 (canceled).

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Claim 30 (new) A method of isolating a nucleic acid molecule associated with a phenotypic trait of interest comprising:

- (A) screening a mapping population of *Arabidopsis* plants to determine the linkage of said phenotypic trait with a collection of polymorphisms, wherein said polymorphisms are distributed throughout the genome of said mapping population of *Arabidopsis* plants at an average density of more than one polymorphism per about 100kb;
- (B) calculating the linkage of each of said polymorphism to said phenotypic trait; and
- (C) isolating said nucleic acid molecule associated with said phenotypic trait based on its linkage to one or more of said polymorphisms.

wherein said collection of polymorphisms is capable of detecting Single Nucleotide Polymorphism 471736.